

Figure 3

5 '		11	21	31	41
	, TGTATGGGA	ATAGTGTTT <u>C</u>	CATATGATCT	GTTGTCTGGA	STATATGCTAC CATATACGATG
•	ACATACCCT	ratcacaaa <u>b</u>	<u>GTATACTAGA</u>	CAACAGACCT	CATATACGATG
 .					probe 1
5 '		61	71	8T	91
5:	1 MACAACHAA7	LACIGIACAA VTCVCVTCTT	AAACCCAGTG	CAGCTGATGAT	rgcaaagcagt Acgtttcgtca
	IACAAGIAA	11GACAIGII	1 1 1 GGG I CAC	GICGACIACIA	ACGITICGICA
5 '		11	21	31	41
	. CTCTCTCTGT	GTACAGTGC	CCCACCTATT	TAAAAATCAC	TACTTGCCCA
103	L GAGAGAGACA	CATGTCACG	GGTGGATAA	ATTTTTAGTGO	TACTTGCCCA CATGAACGGGT
5 '		61	71	81	91
151	GAACACTGTG	BAAACACTTA	A CATAAGAACI	AAACGCAGCGT	CTGGATTCTT GACCTAAGAA
101	CTTGTGACAC	TTTGTGAAT	rgtattettg:	TTTGCGTCGCA	GACCTAAGAA
		7.7 Ombo 2	0.5		
5'		TT brook 5	21	31	41 AAGAGGTCCG TTCTCCAGGC
201	TCCAAGGAGA	GCAGCTTTC	CCACAGGAAC	CACAGTAACAA	AAGAGGTCCG
	AGGIICCICI	CGICGAAAGA	AGGIGICCTIC	FIGICALIGIA	TICICCAGGC
5.1	•	61	71	81	91
051	CCGCCATCCA	CACCCAGCCA	AGACACCTCA	GAGGCCATAG	GGACAACCTC CCTGTTGGAG
251	GGCGGTAGGT	GTGGGTCGGT	'TCTGTGGAGI	CTCCGGTATC	CCTGTTGGAG
5'		11	21	31	41
301	CTTGCTGGCC GAACGACCGG	AACACCTGCT	'GGAGCAGGGG	CACAGGTCCC	AGCAACTGAT
	GAACGACCGG	TTGTGGACGA	CCTCGTCCCC	GTGTCCAGGG	TCGTTGACTA
51		61	71	81	91
					TCTTTTGAAG
351	CCTCAGTGGA GGAGTCACCT	ACCCAGACGT	CGGTTTCGGA	ATTACCCGAG	AGAAAACTTC
5 '		11	21	31	41
401	GGGAAAGAAA CCCTTTCTTT	GAATTTCAAG	CTTATGATAT	CCAATATTAT	TATAGTTGAT
101	CCCTTTCTTT	CTTAAAGTTC	GAATACTATA	GGTTATAATA	ATATCAACTA
5 '		51	71	81	91
5	ር አር ጥጥ አር ጥ አ	_	, <u>,</u> , , , , , , , , , , , , , , , , , , ,	0.1	. · ·
451	GAGTTAGTAAA CTCAATCATT	TAAGGTTTTT	TTTTT	•	



Figure 5

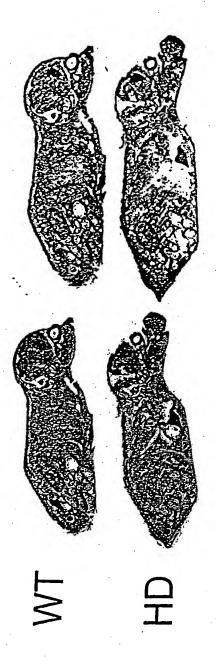
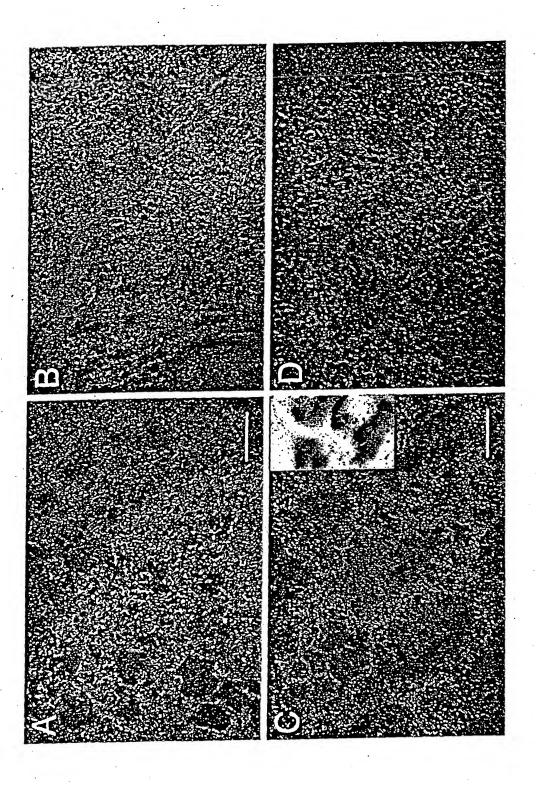
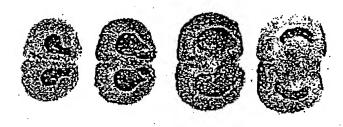


Figure 6





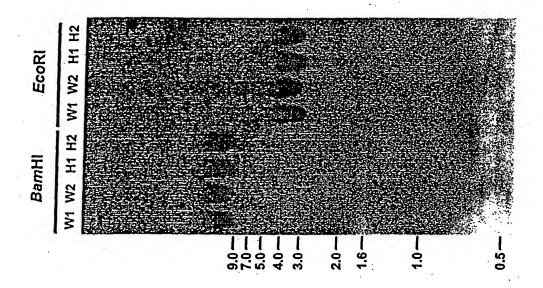
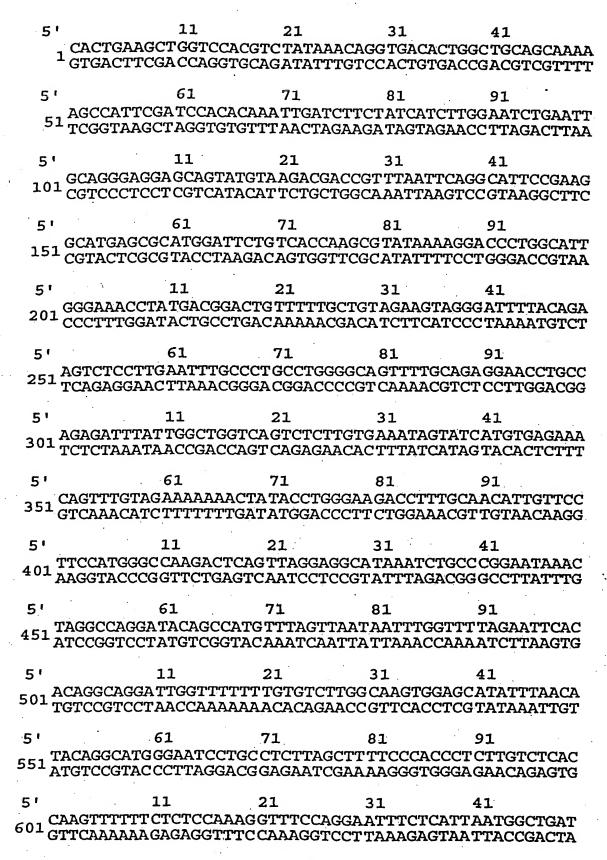
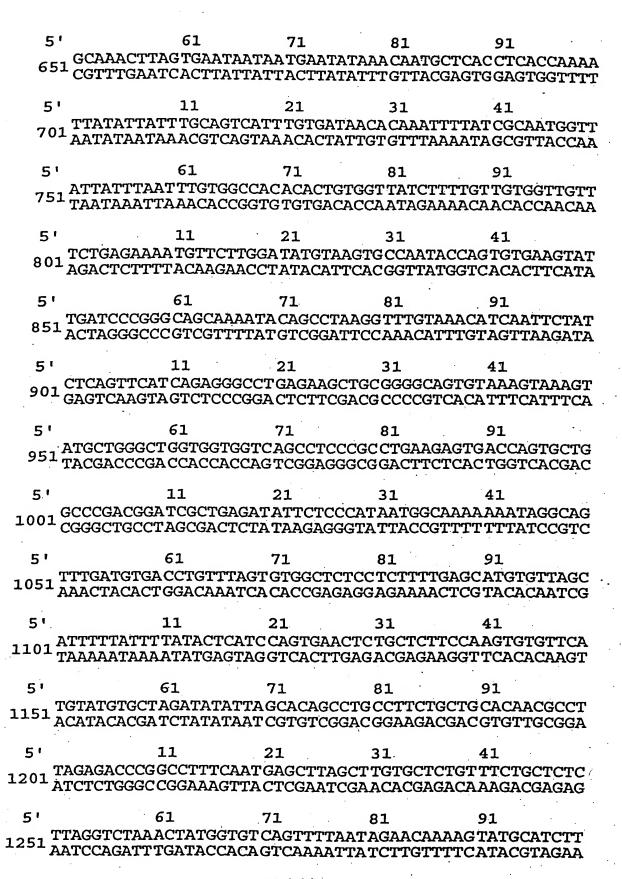
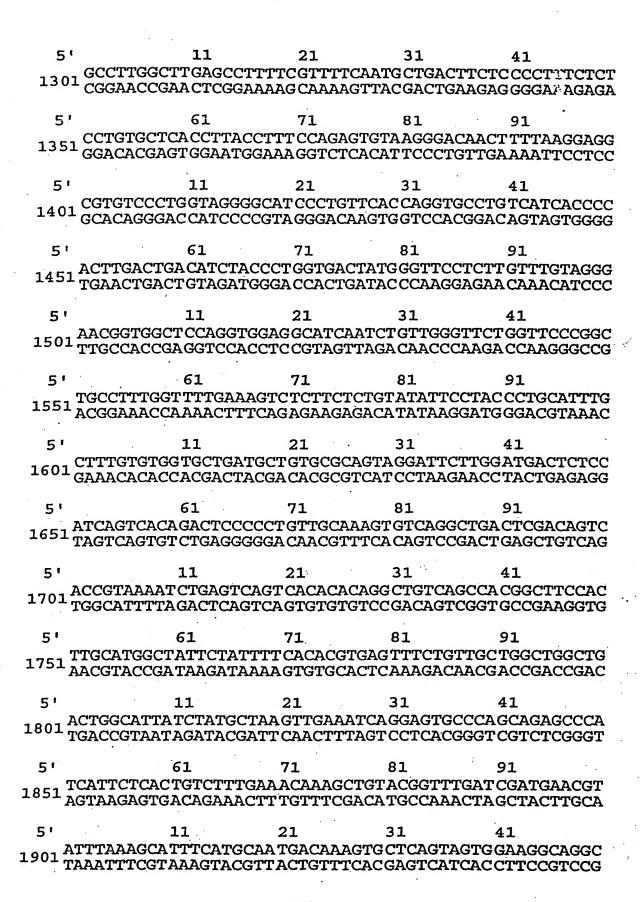
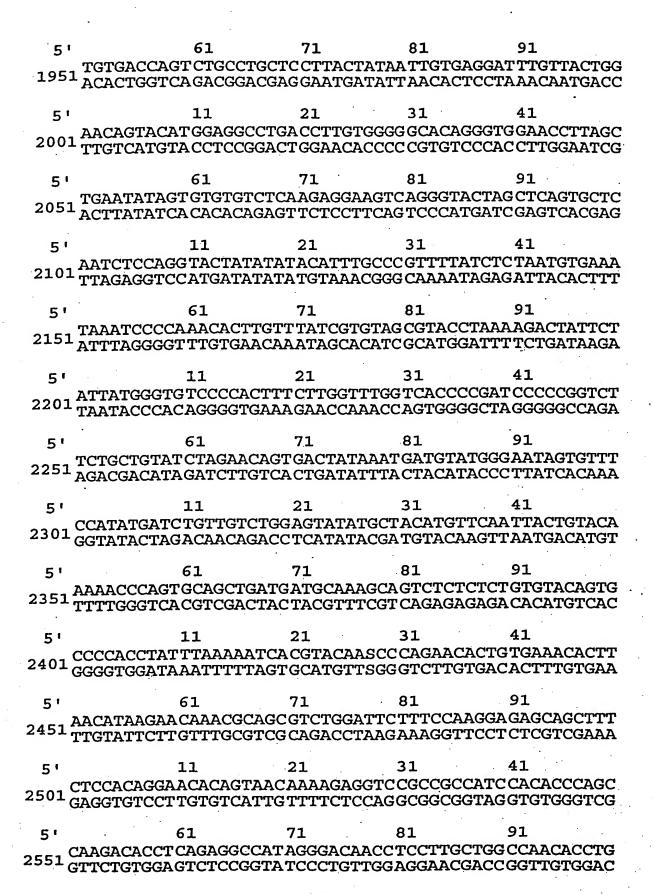


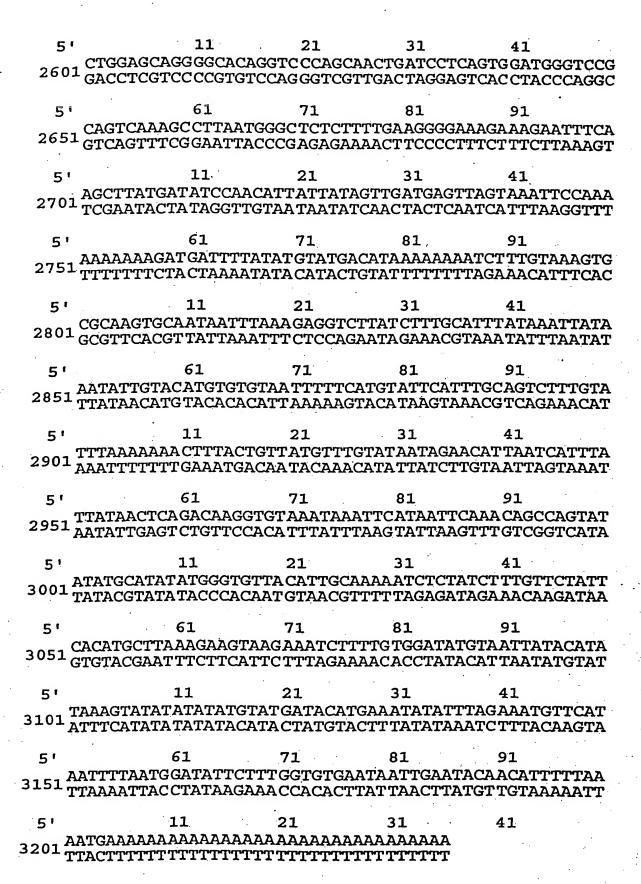
Figure 10











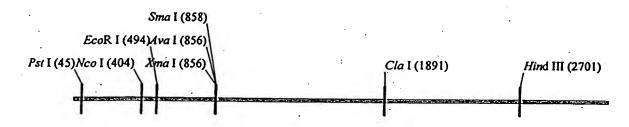
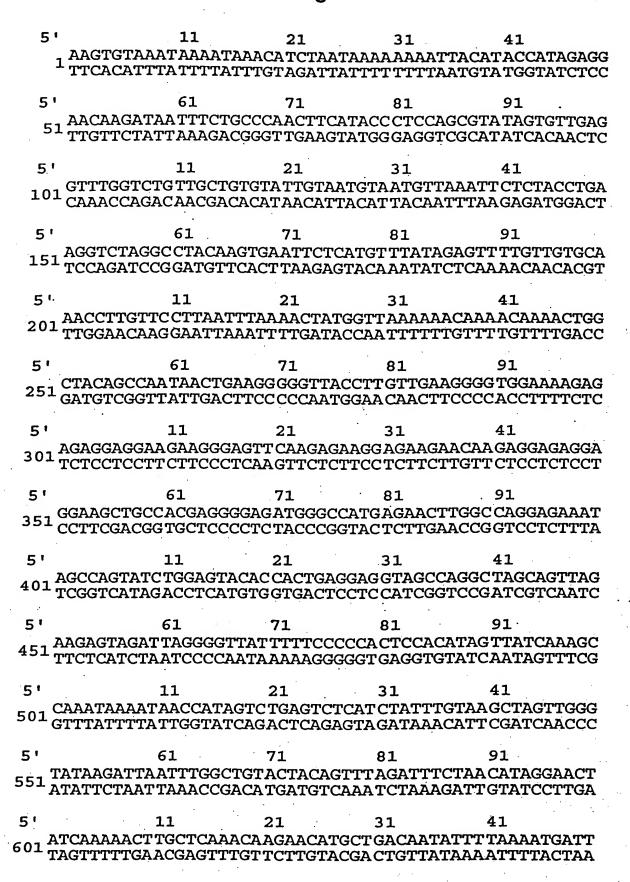
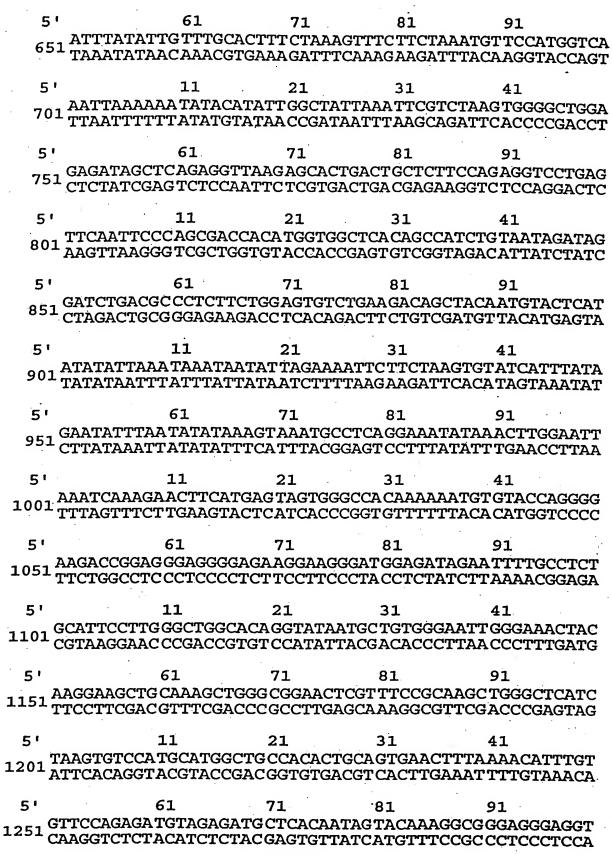
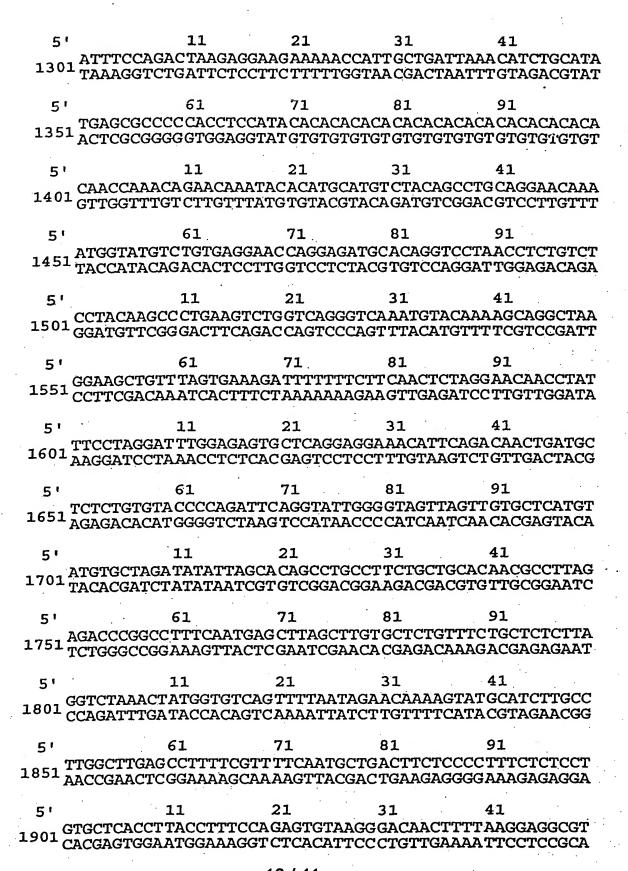


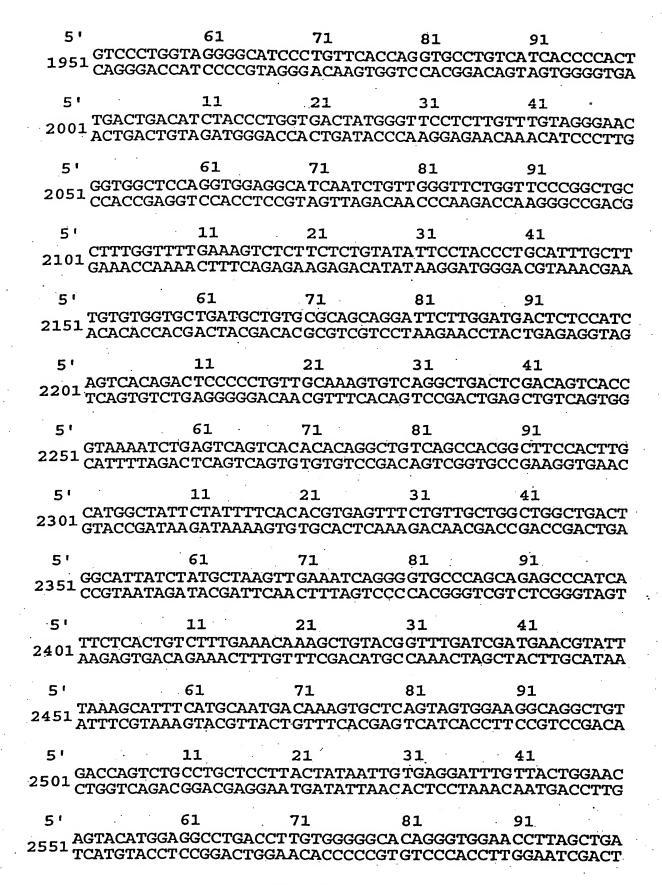
Figure 11 3236 bp

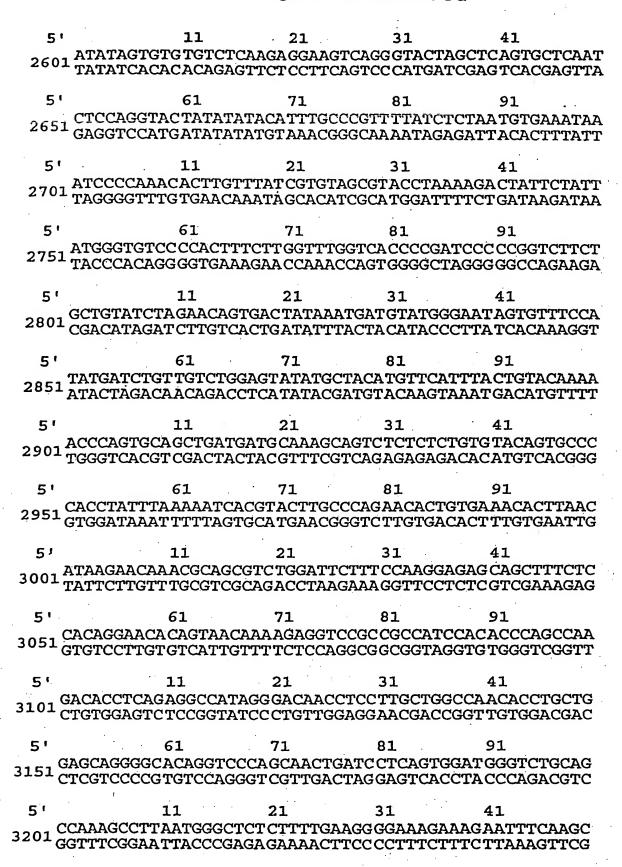
Figure 12

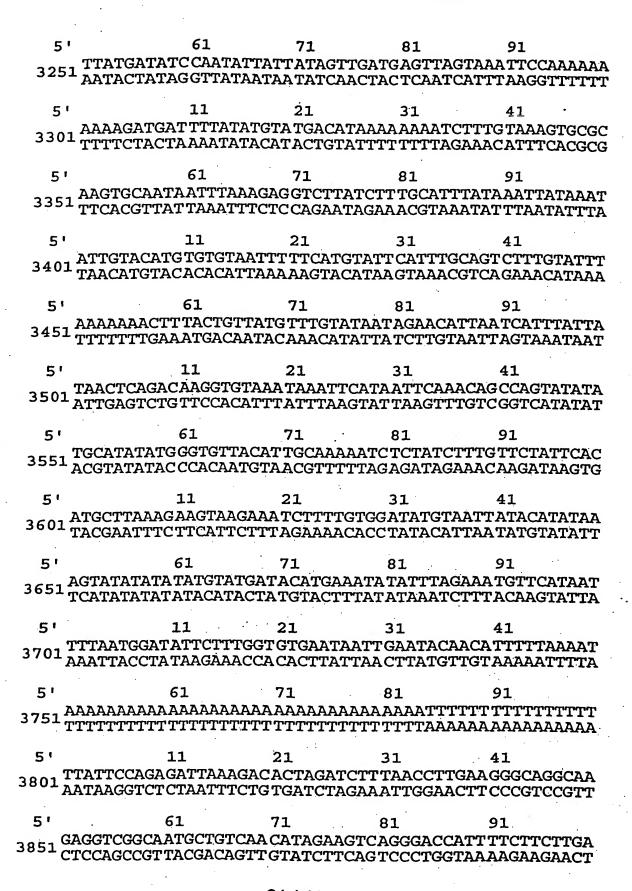


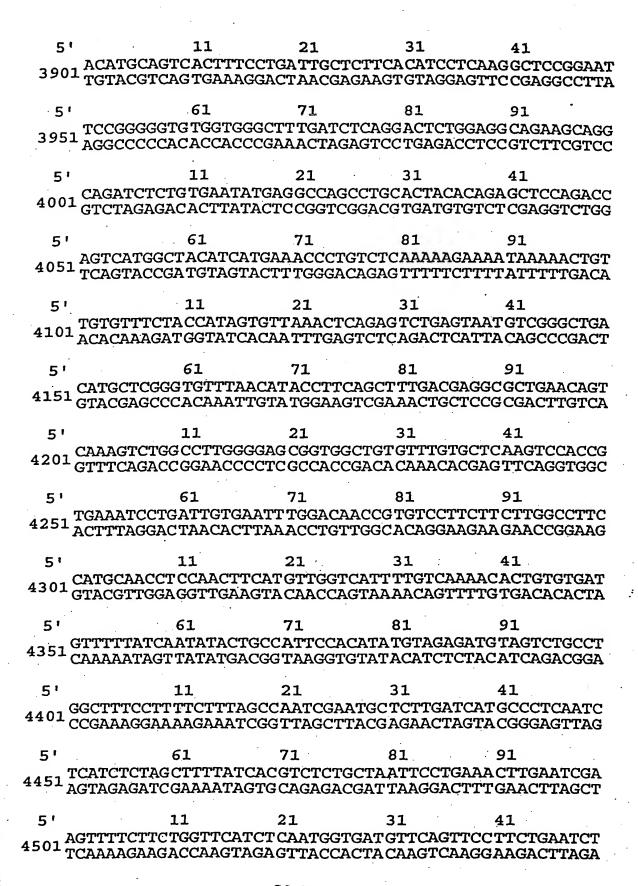


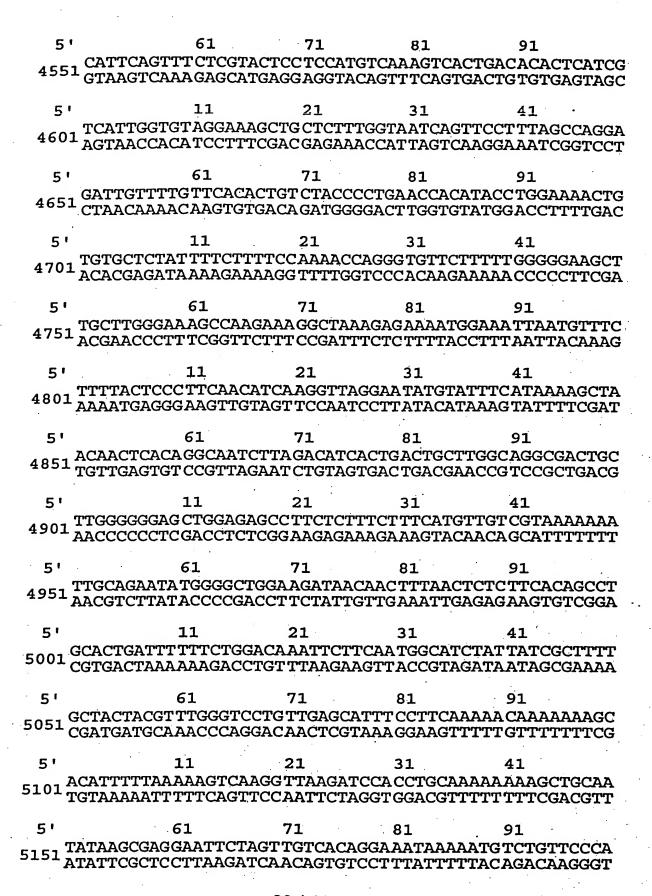


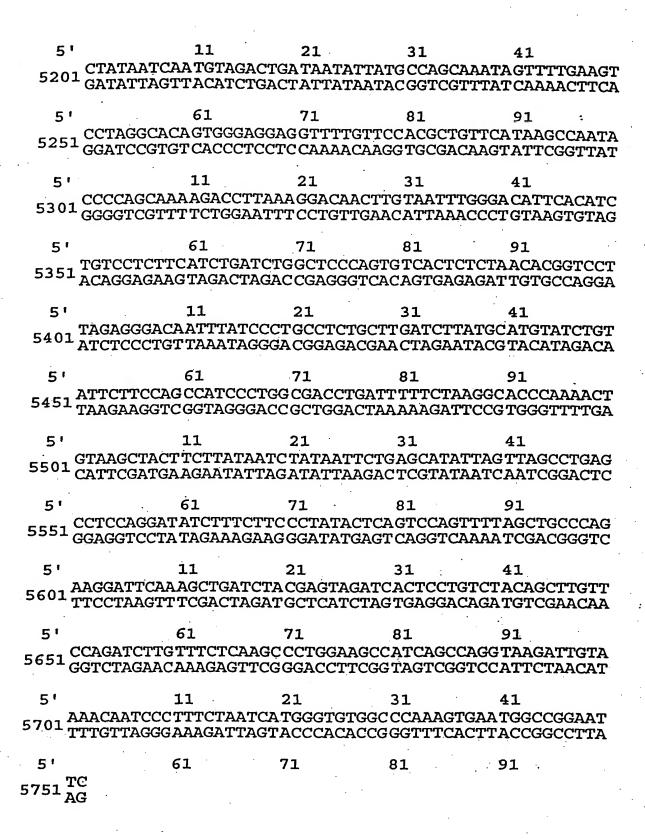


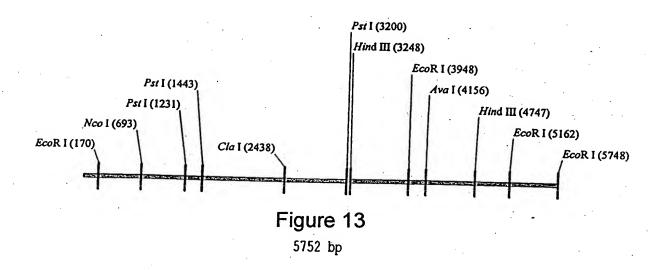


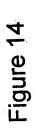












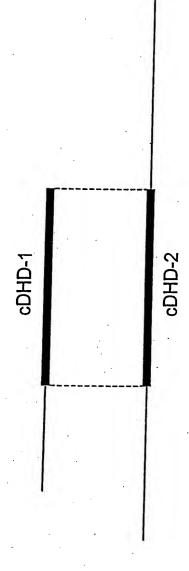


Figure 15

	1	CGCCCGGGCA	GGTCTGTTG	AGGGCAGTTG	GTCAACCTGA	CCAGAGAGAG	CTGAGCTGGA
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•	61			TGCCTTTCAG			
	01	TCTGGGGTGA	CTACCACAC	ACGGAAAGTC	AGGTCCTTCT	TTCTTTCCTT	CCTAAGACTC
	121			CCTGGAGAAG			
		CTAAACCCGT	TTCGGTGTA	GGACCTCTTC	AGACATATGA	CTACGGTTTG	GGTTCTCGAC
	181	AGCTGCTGAT	GAGGCCCAGG	GAGTAGCCCA	CGCGCCCTGA	GCTGTTGGCT	AGCAAGGCCT
	101	TCGACGACTA	CTCCGGGTCC	CTCATCGGGT	GCGCGGGACT	CGACAACCGA	TCGTTCCGGA
·· . ·· ··	241			AAAAATTATA			
	434	AGGACGAGGT	ACACCGTACC	TTTTTAATAT	ACCAAACTGC	CTACTTTTCC	ACTTCCGGAT
	301			TATTAGATGA			
	301	AGAAAGAGAG	GTAGGGGTCC	ATAATCTACT	TAAACAAAGA	CTTTCACAAT	CACGTCTCTG
	361			GGAAAACCAA			
	301	ACACCTTTTC	ACCGACTTCI	CCTTTTGGTT	GTTTCGTTTT	CTACTTGGTA	GAGGGTTCCT
	421	AGTCAGCAGG		CGAATATGCA			
	421	TCAGTCGTCC	ATGGTCCTAT	GCTTATACGT	CCCTCAGCAC	ATGCTCGACT	TGTCGATGTA
	481	AGAGCAGCGC	CTGGACACGG	GCGGGGACAA	CCACCTGCTC	CTCTATGAGC	TCAGCAGCAT
	401	TCTCGTCGCG	GACCTGTGCC	CGCCCCTGTT	GGTGGACGAG	GAGATACTCG	AGTCGTCGTA
	541	CATCAGGATA	GCCACAAAAG	CCGACGGATT	TGCACTGTAC	TTCCTTGGAG	AGTGCAATAA
	341	GTAGTCCTAT	CGGTGTTTTC	GGCTGCCTAA	ACGTGACATG	AAGGAACCTC	TCACGTTATT
	601	TAGCCTGTGT	GTGTTCATAC	CACCCGGGAT	GAAGGAAGGC	CAACCCCGGC	TCATCCCTGC
	001	ATCGGACACA	CACAAGTATG	GTGGGCCCTA	CTTCCTTCCG	GTTGGGGCCG	AGTAGGGACG
• • • •	661	AGGGCCCATC	ACCCAGGGTA	CCACCATCTC	TGCCTACGTG	GCCAAGTCTA	GGAAGACGTT
	001	TCCCGGGTAG	TGGGTCCCAT	GGTGGTAGAG	ACGGATGCAC	CGGTTCAGAT	CCTTCTGCAA
	:		EcoRV		Xhol		
	721	GTTGGTAGAG	GATATCCTTG	GGGATGAGCG	ATTTCCTCGA	GGTACTGGCC	TGGAATCAGG
		CAACCATCTC	CTATAGGAAC	CCCTACTCGC	TAAAGGAGCT	CCATGACCGG	ACCTTAGTCC
	781	AACCCGCATC	CAGTCTGTTC	TTTGCTTGCC	CATTGTCACT	GCCATTGGAG	ACTTGATTGG
		TTGGGCGTAG	GTCAGACAAG	AAACGAACGG	GTAACAGTGA	CGGTAACCTC	TGAACTAACC
	841	CATCCTTGAA	CTGTACAGGC	ACTGGGGCAA	AGAGGCCTTC	TGCCTCAGCC	ATCAGGAGGT
	• • •	GTAGGAACTT	GACATGTCCG	TGACCCCGTT	TCTCCGGAAG	ACGGAGTCGG	TAGTCCTCCA
•	901	TGCAACAGCC	AATCTTGCTT	GGGCTTCCGT		CAGGTGCAGG	
		ACGTTGTCGG	TTAGAACGAA	CCCGAAGGCA	TCGTTATGTG	GTCCACGTCC	ACACATCTCC.
	961	TCTCGCCAAA	CAGACCGAAC	TGAATGACTT	CCTACTCGAC	GTATCAAAGA	CATACTTTGA
	, ,	AGAGCGGTTT	GTCTGGCTTG	ACTTACTGAA	GGATGAGCTG	CATAGTTTCT	GTATGAAACT
• • •	1021	TAACATAGTT	GCCATAGACT	CTCTACTTGA	ACACATCATG	ATATATGCAA	AAAATCTAGT
		ATTGTATCAA	CGGTATCTGA	GAGATGAACT	TGTGTAGTAC	TATATACGTT	TITIAGATCA
	1081	CNACCCCGAC	CCCTCCCCCC	TCTTCCAGGT	GGACCACAAG	AACAAGGAGC	TGTACTCGGA
	100-	CTTGCGGCTG	GCGACGCGCG	AGAAGGTCCA	CCTGGTGTTC	TTGTTCCTCG	ACATGAGCCI
	1141	CCTCTTTCAC	ATTGGGGAGG	AGAAGGAGGG	GAAGCCCATC	TTCAAGAAGA	CCAAGGAGAT
		GGACAAACTG	TAACCCCTCC	TCTTCCTCCC	CTTCGGGTAG	AAGTTCTTCT	GGTTCCTCTA
- • •	1201	CACATTTTCC	ATTGAGAAAG	GGATTGCTGG	TCAAGTGGCA	AGAACAGGCG	AAGTCTTGAA
	•	GTCTAAAAGG	TAACTCTTTC	CCTAACGACC	AGTTCACCGT	TCTTGTCCGC	TTCAGAACTI
	1261	CATTCCCGAT	GCCTACGCGG	ACCCTCGCTT	TAACAGGGAG	GTGGACCTGT	ACACAGGCTA
		GTAAGGGCTA	CGGATGCGCC	TGGGAGCGAA	ATTGTCCCTC	CACCTGGACA	TGTGTCCGAT
	1321	CACCACGAGG	AACATTCTGT	GTATGCCCAT	AGTGAGCCGA	GGCAGCGTGA	TTGGCGTGGT
	·	GTGGTGCTCC	TTGTAAGACA	CATACGGGTA	TCACTCGGCT	CCGTCGCACT	AACCGCACCA

PDE10a and RACEs compiled

	1381	GCAGATGGT	G AACAAGATCA	GCGGTAGCGC	CTTCTCCAAG	ACAGACGAGA	ACAACTTCAA
	1301		C TTGTTCTAGT		GAAGAGGTTC	TGTCTGCTCT	TGTTGAAGTT
			,				BamHI
	3 4 4 3	GATGTTTGC'	r GTCTTCTGCG	CACTGGCCTT	GCACTGTGCT	AACATGTACC	
	1441		A CAGAAGACGC		CGTGACACGA	TTGTACATGG	TGTCCTAGGC
					Hindill	•	
	1501	CCACTCAGAI	A TGCATCTACA	GGGTTACCAT	********	TCCTACCACA	GCATCTGCAC
	1501	CCACTCACAC	r ACGTAGATGT	CCCAATGGTA	CCTCTTCGAA	AGGATGGTGT	CGTAGACGTG
	1.5.61		TGGCAAGGCC			GCACGCATCT	
	1561		ACCGTTCCGG		GTTGGATGGT	CGTGCGTAGA	CGGCCCTGTA
		CGAGCTATTO		TTGGTCCTTT	CGAGAACATG	TGGCCTGGGA	TCTTTGTCTA
	1621		GTGAAACTGT		GCTCTTGTAC	ACCGGACCCT	AGAAACAGAT
		CATGATCCAT		GGACATCCTG		GAAAAATTGT	GCCGTTTTAT
	1681		GCCAGAACAC	CCTGTAGGAC	AAAACTTGAA	CTTTTTAACA	CGGCAAAATA
			AAGAAGAACT			AACTGGAAGC	ATGCAGTCAC
	1741	CMTGTCTGTC	TTCTTCTTGA	TAGCCGCCCA	AGGAATGGTG	TTGACCTTCG	TACGTCAGTG
		GIACAGACAC					Xhol
		GGTGGCACAC	- -	CCATACTTCA	AAACAACAAT	GGCCTCTTCA	
	1801	•••	ACGTACATAC		TTTGTTGTTA	CCGGAGAAGT	GTCTGGAGCT
		Khol	7.001.101.1110			:	· ·
	.061	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	CTGCTAATTG	CGTGTCTGTG	CCATGACCTG	GACCACAGGG	GCTTCAGTAA
	1861	CCCCTTTCCC	GACGATTAAC	GCACAGACAC	GGTACTGGAC	CTGGTGTCCC	CGAAGTCATT
		CACCTACCTG	CAGAAGTTCG	ACCACCCCCT	GGCGGCGCTG	TACTCCACCT	CCACCATGGA
	1921		GTCTTCAAGC		CCGCCGCGAC	ATGAGGTGGA	GGTGGTACCT
	7.001	GCAACACCAC			CCTTCAGCTG	GAAGGCACA	ATATCTTCTC
	1981		AAGAGGGTCT		GGAAGTCGAC	CTTCCCGTGT	TATAGAAGAG
		CACCCTGAGO		ACGAGCAGGT		ATCCGCAAAG	CCATCATCGC
	2041	GTGGGACTCG			CGACCTCTAG	TAGGCGTTTC	GGTAGTAGCG
	01.01	CACCGACCTC			GAAGCAGTTG	GAGGAGATGT	ACCAGACAGG
	2101	GTGGCTGGAG	CGGGATATGA			CTCCTCTACA	TGGTCTGTCC
	21.61	GTCGCTGAAC		AGTCCCATCG	AGACCGTGTC		TGATGACTGC
	2161	CAGCGACTTG	: _ <u>-</u>	TCAGGGTAGC	TCTGGCACAG	TAGCCGAACT	ACTACTGACG
	2221	CTGTGATCTT		CCAAACTATG			CGAATGATAT
	2221	CACACTAGAA	ACGAGACACT	GGTTTGATAC	CGGTCAATGT	TTTAACTGTC	GCTTACTATA
		F	co81		,	:	•
	2201	AADTCCTA	TTCTGGGCTG	AGGGTGATGA	GATGAAGAAG	CTGGGCATAC	AGCCCATTCC
	2281	TATACGTCTT	AAGACCCGAC	TCCCACTACT	CTACTTCTTC	GACCCGIAIG	10000111100
	2241		ACACACAAGC	CAGATGAAGT	CCCTCAAGGG	CAGCTCGGAT	TCTACAATGC
	2341	አምአርጥጀርርጥG	TCTCTGTTCG	CTCTACTTCA	GGGAGTTCCC	GICGMGCCIM	AGAIGIIMGG
	2401		CCCTCCTATA	CCACCTTGAC	GCAGATCCTC	CCACCCACAG	AGCCTCTGCT
	2401	ACACCGGTAA	GGGACGATAT	GGTGGAACTG	CGTCTAGGAG	GGTGGGTGTC	1CGGAGAGA
	2461		ACCCA TRADCC	TCAATCAGTG	GGAGAAGGTA	ATTCGCGGGG	AAGAGACAGC
	2461	CTTCCGGACG	TCCCTATTGG	AGTTAGTCAC	CCTCTTCCAT	TAAGCGCCCC	1101010100
	2521		TCACCCCCAG	GCCCGGCGCC	TAGCAAGAGC	ACACCTGAGA	AGCTGAACGT
	2521	ጥጥአ C D C C T A A	AGTCCGGGTC	CGGGCCGCGG	ATCGTTCTCG	TGTGGACICI	TCGACTTGC:
	25.01	CARCOMMEN A	CACTCATCCT	GAAGTGACGT	CCTGATGTCT	GÇCCAGCAAC	CGACTCAACC
	2581	CTTCCAACTT	CTGACTAGGA	CTTCACTGCA	GGACTACAGA	CGGGTCGTTG	GCIGAGIICO
	2641	#COMMOMCMC	A CTTCCTTCT	TTTTGTTTTC	AAGGGGTGAA	AACCCCCTGT	CAGAAGGTAC
	2641	ACGAAGACAC	TGAAGCAAGA	AAAACAAAAG	TTCCCCACTT	TTGGGGGACA	GTCTTCCATG
				·		•	•

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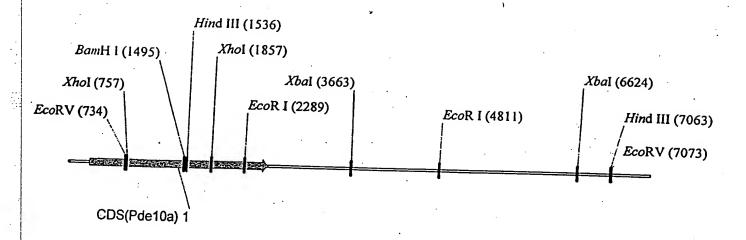
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	2761	CCAACCCAGG CTCTGCCGTG TTCAGACGTC	GGCTACTCCG TGGCTCCACC TGACCTCCGA
	2701	CGTTGGGTCC GAGACGGCAC AAGTCTGCAG	
	2821	ATGCTATTTG CTCCCAGGCC AGCACTGCAC	
	2021	TACGATAAAC GAGGGTCCGG TCGTGACGTG	ACAGACCTCC CCCGTCTCTG GTGTCCTCTC
	2001	GTTCTTGCCT GCATCCTCCC ATGAGGGTGT	
	2881	CAGRACGGA CGTAGGAGGG TACTCCCACA	CCGGTCAAGG GATCAAGACA CGGTACGACG
		TCCTTGGTGG CATTGGTTAG GAATGGGACA	CACGCCCTT GTTGTGAAGT TTACATGTGA
	2941	ACGARCACC GTAACCAATC CTTACCCTGT	GTGCGGGGAA CAACACTTCA AATGTACACT
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	3001	GGAGAATAT CCAATTGACT CAAACACCGG	ACCTGTGTAC ATTACTTCCA GTGTCAGGTG
		ACCUCACACA CARATCCARA CTGTTGATTA	CAGGTGCACT ACAGGTATGC TCTTTCAGTC
	3061	TCCACTGTCT CTTTAGGTTT GACAACTAAT	GTCCACGTGA TGTCCATACG AGAAAGTCAG
		TRUTCECCC CACATAGGTG AGTCTGCTCC	ACTCAGAANN AAGCATACCT CTGCCCTCAT
	3121	ATACACCCC GTGTATCCAC TCAGACGAGG	TGAGTCTTNN TTCGTATGGA GACGGGAGTA
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<u>.</u>		TORRODATE ARROCACTC CCCTCCCCT	CACTGTAGCC TTCGACAACT GCGCCAATCC
	3241	ACTUTOTIAN TITTGTGGAG GGGAGGGGGA	GTGACATCGG AAGCTGTTGA CGCGGTTAGG
		COTTO TACAL ACADALTARA ACTARGCAT	ATAAATTTCC TCCAGCAAGC AAATCTTGTG
	3301	GAAATATGTT TCTTTTATTT TCATTCCGTA	TATTTAAAGG AGGTCGTTCG TTTAGAACAC
		CCTARARARA ARCCATGTGA ATNNTAACAA	CNTCTANANT NTCNCNGNAT GTTATGGCAG
	3361	CCATTTTTT TTCGTACACT TANNATTGTT	GNAGATNINA NAGNGNCNIA CAATACCGIC
		AATTTTAGTC ACGTCCAAAA CAAAAAGATT	ATTCCAGAAG ATACCTCATC CTATGCCTGA
	3421	TTAAAATCAG TGCAGGTTTT GTTTTTCTAA	TAAGGTCTTC TATGGAGTAG GATACGGACT
	2403	PACCETCEAC AGENTGGCGT CCGTCTCCCA	GGGTTCTGAT CCGTCTCCTC ACGGTGCAAT
	3481	TTCCGAGGTG TCGTACCGCA GGCAGAGGGT	CCCAAGACTA GGCAGAGGAG TGCCACGTTA
	2541	CACCAGGAC AGAGAGGAGG GCTGCAGGGC	TACCACATTG ACCCAGAAGG TATCTCCTCT
	3541	GTCCGTCCTG TCTCTCCTCC CGACGTCCCG	ATGGTGTAAC TGGGTCTTCC ATAGAGGAGA
	2601	CACCATTCAC ACATCCATAA GGAATGCCAA	ATGCTGTATT GAATAGTTCT CTGTGTGACT
	3601	CTGGTAAGTC TGTAGGTATT CCTTACGGTT	TACGACATAA CTTATCAAGA GACACACTGA
		Xhal	
	2662	TTCTAGAGAA GCCAGGACAC CCTGAGCCTT	TCCNGGGGAA CTCTAAGGAG TCACAGGTTC
	3661	AACATCTCTT CGGTCCTGTG GGACTCGGAA	AGGNCCCCTT GAGATTCCTC AGIGICCAAG
	2721	PORCECCE CATTTTCAGE ATAGCATGGA	GACAGAGATC CGGTCGTTGT TCTCACTCGT
	3721	TOTAL COLOR CTARANGTCC TATCGTACCT	CTGTCTCTAG GCCAGCAACA AGAGTGAGCA
		TOTAL ANCENCEDE TEACCAGAAA	CACTCACTCA GCACTCTGCA GGAGCAGGAG
	3781	CTCGGAACTC TTCCTCTCTG ACTGGTCTTT	GTGAGTGAGT CGTGAGACGT CCTCGTCCTC
		TOTAL COME ANCATCA ATC TTGGATAGAT	TTTGATACAC CCAATACCAT ACACACAGGA
	3841	TTCTATGARA TTCTACTTAG AACCTATCTA	AAACTATGTG GGTTATGGTA TGTGTGTCCT
	2001	COMMCCOATT TOCALACTOT ATTOAGTTTC	CTTCCGCGCT CTGACCCACG GTTGTAGCGG
	3901	CCAACCGTAA ACGTTTCAGA TAAGTCAAAG	GAAGGCGCGA GACTGGGTGC CAACATCGCC
	2061	ACTOR ACACTOTARC ACTOTACATO	CGATTTCCCC ATGGGCTTCT AAAATGTCAC
	3961	TCACCCGACT TGTGACATTG TGACATGTAC	GCTAAAGGGG TACCCGAAGA TITTACAGTG
	4021	CARCTCCTCC CCTGCTGTGT CCTACTCCAT	TTACTGGTTA CAAGGTGATG TCAACAAGAG
	4021	GTAGAGGAGG GGACGACACA GGATGAGGTA	AATGACCAAT GTTCCACTAC AGTTGTTCTC

						•	
	4081	AAGCTATCAC	AACACCAGGG	CTGTGCACAC	GTGCACACAC	ATGTATGCAC	AAGCACACAG
					CACGTGTGTG		
	4141	ATGTATGTAC	AGCACACACA	CACACACACA	CCCCAAAAGG	AGAGAAAAGG	AAGAAAACAT
	• • • • • • • • • • • • • • • • • • • •	TACATACATG	TCGTGTGTGT	GTGTGTGTGT	GGGGTTTTCC	TCTCTTTTCC	TTCTTTTGTA
	4201	TTATAAAAAG	CGACAGCTAC	CCCATATCAA	AATAGTCTTT	CCTGTAGGAA	ACAGGAGCTC
		AATATTTTC			TTATCAGAAA	GGACATCCTT	TGTCCTCGAG
•	4261			GTGTGTTCTC	CCATCAGTGC	ACTCTCCCAG	GGGTGCTCAC
	1202	AGGTATTCCT		CACACAAGAG	GGTAGTCACG	TGAGAGGGTC	CCCACGAGTG
	4321	TGAAGCTGGT	CCACRTCTAT	AAACAGGTGA	CACTGGCTGC	AGCAAAAAGC	CATTCGATCC
	7,721	ACTTCGACCA	GGTGRAGATA	TTTGTCCACT	GTGACCGACG	TCGTTTTTCG	GTAAGCTAGG
	4381	ACACAAATTG	ATCTTCTATC	ATCTTGGAAT	CTGAATTGCA	GGGAGGAGCA	GYATGTAAGA
	1501	TGTGTTTAAC	TAGAAGATAG	TAGAACCTTA	GACTTAACGT	CCCTCCTCGT	CYTACATTCT
	4441	CGACCGTTTA	ATTCAGGCAT	TCCGAAGGCA	TGAGCGCATG	GATTCTRTCA	CCAAGCGTAT
	3	GCTGGCAAAT	TAAGTCCGTA	AGGCTTCCGT	ACTCGCGTAC	CTAAGARAGT	GGTTCGCATA
	 4501				CGGACTGTTT		
	1301	TTTTCCTGGG	ACCGTAACCC	TTTGGATACT	GCCTGACAAA	AACGACATCT	TCATCCCTAA
	4561				TGGGGCAGTT	TTGCAGAGGA	ACCTGCCAGA
	1001	AATGTCTTCA	GAGGAACRTA	AACGGGACGG	ACCCCGTCAA	AACGTCTCCT	TGGACGGTCT
	4621	CATTTATTGG	CTGGTCAGTC	TCTTGTGAAA	TAGTATCATG	TGAGAAACAG	TTTGTAGAAA
		CTAAATAACC	GACCAGTCAG	AGAACACTTT	ATCATAGTAC	ACTCTTTGTC	AAACATCTTT
	4681	AAAACTATAC	CTGGGAAGAC	CTTTGCAACA	TTGTTCCTTC	CATGGGCCAA	GACTCAGTTA
		TTTTGATATG	GACCCTTCTG	GAAACGTTGT	AACAAGGAAG	GTACCCGGTT	CTGAGTCAAT
•	4741	GGAGGCATAA	ATCTGCCCGG	AATAAACTAG	GCCAGGATAC	AGCCATGTTT	AGTTAATAAT
		CCTCCGTATT	TAGACGGGCC	TTATTTGATC	CGGTCCTATG	TCGGTACAAA	TCAATTATTA
			EcoRI			•	
	4801	ттссттттАС	AATTCACACA	GGCAGGATTG	GTTTTTTGT	GTCTTGGCAA	GTGGAGCATA
		AACCAAAATC	TTAAGTGTGT	CCGTCCTAAC	CAAAAAAACA	CAGAACCGTT	CACCTCGTAT
	4861	TTTAACATAC	AGGCATGGGA	ATCCTGCCTC	TTAGCTTTTC	CCACCCTCTT	GTCTCACCAA
		AAATTGTATG	TCCGTACCCT	TAGGACGGAG	AATCGAAAAG	GGTGGGAGAA	CAGAGTGGTT
	4921	GTTTTTTCTC	TCCAAAGGTT	TCCAGGAATT	TCTCATTAAT	GGCTGATGCA	AACTTAGTGA
		CAAAAAAGAG	AGGTTTCCAA	AGGTCCTTAA	AGAGTAATTA	CCGACTACGT	TIGARICACI
•	4981	ATAATAATGA	ATATAAACAA	TGCTCACCTC	ACCAAAATTA	TATTATTTGC	AGICATIIGI TCACTAAACA
		TATTATTACT				ATAATAAACG	COCOCCOON
	5041	GATAACACAA	ATTTTATCGC	AATGGTTATT		TGGCCACACA	
		CTATTGTGTT	TAAAATAGCG	TTACCAATAA	TAAATTAAAC	ACCGGTGTGT	A MA COA CECT
	5101	CTTTTGTTGT	GGTTGTTTCT	GAGAAAATGT	TCTTGGATAT	GTAAGTGCCA CAMBCACCCT	TATECTCACA
•		GAAAACAACA	CCAACAAAGA	CTCTTTTACA	AGAACCTATA	CATICACOGI	A DECENTATION OF THE PARTY OF T
	5161	GAAGTATTGA	TCCCGGGCAG	CAAAATACAG	CCTAAGGTTT	GTAAACATCA	TARCATACAC
		CTTCATAACT	AGGGCCCGTC	GTTTTATGTC	GGATTCCAAA	CATTIGIAGI	CMCCCCMCCT
	5221	AGTTCATCAG	AGGGCCTGAG	AAGCTGCGGG	GCAGTGTAAA	GTAAAGTATG	CACCCGACCA
		TCAAGTAGTC	TCCCGGACTC	TTCGACGCCC	CGTCACATTT	CATTICATAC	CCTCCCTCCA
	5281	GGTGGTCAGC	CTCCCCTTGC	CAAGAAGAGA	GCAATTGAAT	CCTGTCCCCA	CCACCGACCT
		CCACCAGTCG	GAGGGGAACG	GTTCTTCTCT	CGTTAACTTA	CARA MECECCI	CATABTECCA
		CGCCTGAAGA	GTGACCAGTG	CTGGCCCGAC	GGATCGCTGA	CONTRACTOR	CHARTIGGON
	· .	GCGGACTTCT	CACTGGTCAC	GACCGGGCTG	CCTAGCGACT	CIMINAGAGG	ACCATE TT
	5401	AAAAAATAGG	CAGTTTGATG	TGACCTGTTT	AGTGTGGCTC	ACCACAAAA	TCGTACACAA
		TTTTTTATCC	GTCAAACTAC	ACTGGACAAA	TCACACCGAG	AGGAGAAAAC	10011101101

	5461	AGCATTTTTA	TTTTATACTC	ATCCAGTGAA	CTCTGCTCTT	CCAAGTGTGT	TCATGTATGT
	3.01	TCGTAAAAAT	AAAATATGAG	TAGGTCACTT	GAGACGAGAA	GGTTCACACA	AGTACATACA
٠	5521	CCTAGATATA	TTAGCACAGC	CTGCCTTCTG	CTGCACAACG	CCTTAGAGAC	CCGGCCTTTC
	3321	CGATCTATAT	AATCGTGTCG	GACGGAAGAC	GACGTGTTGC	GGAATCTCTG	GGCCGGAAAG
•	5581	AATGAGCTTA	GCTTGTGCTC	TGTTTCTGCT	CTCTTAGGTC	TAAACTATGG	TGTCAGTTTT
	3301	TTACTCGAAT	CGAACACGAG	ACAAAGACGA	GAGAATCCAG	ATTTGATACC	ACAGTCAAAA
	5641	AATAGAACAA	AAGTATGCAT	CTTGCCTTGG	CTTGAGCCTT	TTCGTTTTCA	ATGCTGACTT
	3041	TTATCTTGTT	TTCATACGTA	GAACGGAACC	GAACTCGGAA	AAGCAAAAGT	TACGACTGAA
	5701	CTCCCCTTTC	TCTCCTGTGC	TCACCTTACC	TTTCCAGAGT	GTAAGGGACA	ACTTTTAAGG
	, 3701	GAGGGGAAAG	AGAGGACACG	AGTGGAATGG	AAAGGTCTCA	CATTCCCTGT	TGAAAATTCC
	5761	AGGCGTGTCC	CTGGTAGGGG	CATCCCTGTT	CACCAGGTGC	CTGTCATCAC	CCCACTTGAC
	5701	TCCGCACAGG	GACCATCCCC	GTAGGGACAA	GTGGTCCACG	GACAGTAGTG	GGGTGAACTG
	5821	TGACATCTAC	CCTGGTGACT	ATGGGTTCCT	CTTGTTTGTA	GGGAACGGTG	GCTCCAGGTG
	3021	ACTGTAGATG	GGACCACTGA	TACCCAAGGA	GAACAAACAT	CCCTTGCCAC	CGAGGTCCAC
	5881	GAGGCATCAA	TCTGTTGGGT	TCTGGTTCCC	GGCTGCCTTT	GGTTTTGAAA	GTCTCTTCTC
	3661	CTCCGTAGTT	AGACAACCCA	AGACCAAGGG	CCGACGGAAA	CCAAAACTTT	CAGAGAAGAG
	5941	TGTATATTCC	TACCCTGCAT	TTGCTTTGTG	TGGTGCTGAT	GCTGTGGCAG	TAGGATCTTG
	7341	ACATATAAGG	ATGGGACGTA	AACGAAACAC	ACCACGACTA	CGACACCGTC	ATCCTAGAAC
	6001	CATGACTCTC	CATCAGTCAC	AGACTCCCCC	TGTTGCAAAG	TGTCAGGCTG	ACTCGACAGT
	0001	CTACTGAGAG	GTAGTCAGTG	TCTGAGGGGG	ACAACGTTTC	ACAGTCCGAC	TGAGCTGTCA
	6061	CACCGTAAAA	TCTGAGTCAG	TCACACACAG	GCTGTCAGCC	ACGGCTTCCA	CTTGCATGGC
	0001	GTGGCATTTT	AGACTCAGTC	AGTGTGTGTC	CGACAGTCGG	TGCCGAAGGT	GAACGTACCG
	6121	ΤΑ ΤΤΟ ΤΑ ΤΤΤ	TCACACGTGA	GTTTCTGTTG	CTGGCTGGCT	GACTGGCATT	ATCTATGCTA
	0121	ATAAGATAAA	AGTGTGCACT	CAAAGACAAC	GACCGACCGA	CTGACCGTAA	TAGATACGAT
	6181	AGTTGAAATC	AGGAGTGTGC	CCAGCAGAGC	CCATCATTCT	CACTGTCTTT	GAAACAAAGC
	0	TCAACTTTAG	TCCTCACACG	GGTCGTCTCG	GGTAGTAAGA	GTGACAGAAA	CTTTGTTTCG
	6241	TGTACGGTTT	GATCGATGAA	CGTATTTAAA	GCATTTCATG	CAATGACAAA	GTGCTCAGTA
		ACATGCCAAA	CTAGCTACTT	GCATAAATTT	CGTAAAGTAC	GTTACTGTTT	CACGAGTCAT
	6301	GTGGAAGGCA	GGCTGTGACC	AGTCTGCCTG	CTCCTTACTA	TAATTGTGAG	GATTTGTTAC
		CACCTTCCGT	CCGACACTGG	TCAGACGGAC	GAGGAATGAT	ATTAACACTC	CTAAACAATG
	6361	TGGAACAGTA	CATGGAGGCC	TGACCTTGTG	GGGCACAGG	GTGGAACCTT	AGCTGAATAT
		ACCTTGTCAT	GTACCTCCGG	ACTGGAACAC	CCCCGTGTCC	CACCTTGGAA	TCGACTTATA
	6421	AGTGTGTGTC	TCAAGAGGAA	GTCAGGGTAC	TAGCTCAGTG	CTCAATCTCC	AGGTACTATA
		TCACACACAG	AGTTCTCCTT	CAGTCCCATG	ATCGAGTCAC	GAGTTAGAGG	TCCATGATAT
	6481	TATACATTTG	CCCGTTTTAT	CTCTAATGTG	AAATAAATCC	CCAAACACTT	GTTTATCGTG
		ATATGTAAAC	GGGCAAAATA	GAGATTACAC	TTTATTTAGG	GGTTTGTGAA	CAAATAGCAC
•	6541	TAGCGTACCT	AAAAGACTAT	TCTATTATGG	GTGTCCCCAC	TTTCTTGGTT	TGGTCACCCC
		ATCGCATGGA	TTTTCTGATA	AGATAATACC	CACAGGGGTG	AAAGAACCAA	ACCAGTGGGG
	•			Xbal			
	6601	GATCCCCCGG	TCTTCTGCTG	TATCTAGAAC	AGTGACTATA	AATGATGTAT	GGGAATAGTG
	•	CTAGGGGGCC	AGAAGACGAC	ATAGATCTTG	TCACTGATAT	TTACTACATA	COLIMICAC
	6661	TTTCCATATG	ATCTGTTGTC	TGGAGTATAT	GCTACATGTT	CATTTACTGT	ACAAAAACCC
		AAAGGTATAC '	TAGACAACAG	ACCTCATATA	CGATGTACAA	GTAAATGACA	1611111666
-	6721	AGTGCAGCTG	ATGATGCAAA	GCAGTCTCTC	TCTGTGTACA	GTGCCCCACC	ፕ <u>ለ</u> ፓፒፕለለለለለ አመአ አ አጥጥጥጥ
		TCACGTCGAC	TACTACGTTT	CGTCAGAGAG	AGACACATGT	CACGGGTGG	WINNUTITY
	6781	TCACGTACAA	NCCCAGAACA	CTGTGAAACA	CTTAACATAA	GAAACAAACG	CAGCGTCTGG
		AGTGCATGTT	NGGGTCTTGT	GACACTTTGT	GAATTGTATT	CITIGITIEC	0100010100
		•					

6841	ATTCTTTCC	A AGGAGAGCAG	CTTTCTCCAC	AGGAACACAG	TAACAAAAGA	GGTCCGCCGC
	TAAGAAAGG	T TCCTCTCGTC	GAAAGAGGTG	TCCTTGTGTC	ATTGTTTTCT	CCAGGCGGCG
6901	CATCCACAC	CAGCCAAGAC	ACCTCAGAGG	CCATAGGGAC	AACCTCCTTG	CTGGCCAACA
	GTAGGTGTG	GTCGGTTCTG	TGGAGTCTCC	GGTATCCCTG	TTGGAGGAAC	GACCGGTTGT
6961	CCTGCTGGAC	CAGGGCACAG	GTCCCAGCAA	CTGATCCTCA	GTGGATGGGT	CCGCAGTCAA
	GGACGACCTC	GTCCCGTGTC	CAGGGTCGTT	GACTAGGAGT	CACCTACCCA	GGCGTCAGTT
					HindIII	EcoRV
7021	AGCCTTAATG	GGCTCTCTTT	TGAAGGGGAA	AGAAANNTTT		ATATCCAACA
	TCGGAATTAC	CCGAGAGAAA	ACTTCCCCTT	TCTTTNNAAA	GTTCGAATAC	TATAGGTTGT
7081	TTATTATAGT	TGATGAGTTA	GTAAATTCCG	AAAAAAAAG	ATGATTTAT	ATGTATGACA
•	AATAATATCA	ACTACTCAAT	CATTTAAGGC	TTTTTTTTC	TACTAAAATA	
7141	TAAAAAAAT	CTTTGTAAAG	TGCGCAAGTG	CAATAATTTA	AAGAGGTCTT	ATCTTTGCAT
	- ATTTTTTTA	GAAACATTTC	ACGCGTTCAC	GTTATTAAAT	TTCTCCAGAA	TAGAAACGTA
7201	TTATAAATTA	TAAATATTGT	ACATGTGTGT	AATTTTTCAT	GTATTCATTT	GCAGTCTTTG
	AATATTTAAT	ATTTATAACA	TGTACACACA	TTAAAAAGTA	CATAAGTAAA	CGTCAGAAAC
7261	TATTTAAAAA	AACTTTACTG	TTATGTTTGT	ATAATAGAAC	ATTAATCATT	TATTATAACT
	ATAAATTTTT	TTGAAATGAC	AATACAAACA	TATTATCTTG	TAATTAGTAA	ATAATATTGA
7321	CAGACAAGGT	GTAAATAAAT	TCATAATTCA	AACAGCCAGT	ATATATGCAT	ATATGGGTGT
		CATTTATTTA				
7381	TACATTGCAA	AAATCTCTAT	CTTTGTTCTA	TTCACATGCT	TAAAGAAGTA	AGAAATCTTT
	ATGTAACGTT	TTTAGAGATA	GAAACAAGAT	AAGTGTACGA	ATTTCTTCAT	TCTTTAGAAA
7441	TGTGGATATG	TAATTATACA	TATAAAGTAT	ATATATATGT	ATGATACATG	AAATATATTT
	ACACCTATAC	ATTAATATGT	ATATTTCATA	TATATATACA	TACTATGTAC	TTTATATAAA
7501	AGAAATGTTC	ATAATTTTAA	TGGATATTCT	TTGGTGTGAA	TAATTGAATA	CAACATTTTT
	TCTTTACAAG	TATTAAAATT	ACCTATAAGA	AACCACACTT	ATTAACTTAT	GTTGTAAAAA
7561	AAAATGAAAA	AAAAAAAAA	C			
	TTTTACTTTT	TTTTTTTTT	G .		•	
• •				• •		• • • • • • • • • • • • • • • • • • • •

Figure 16



PDE10a and RACEs compiled 7581 bp

Figure 17

PDE10A compiled - coding sequence and features

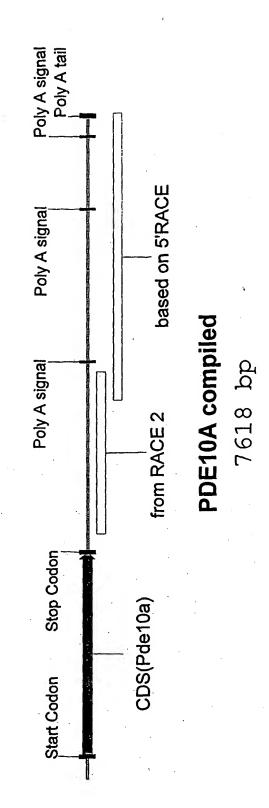
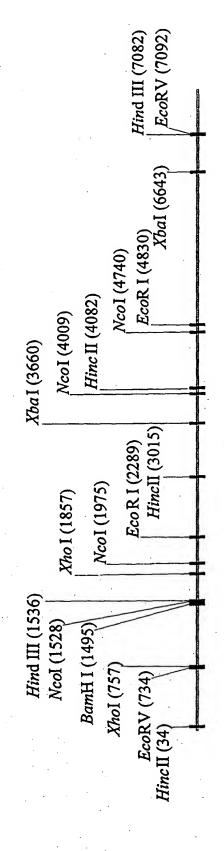


Figure 18

PDE10A compiled - restriction sites



PDE10A compiled 7618 bp

Figure 19

•	1	CGCCCGGGC	A GGTCTGTTGG	AGGGCAGTT	G GTCAACCTGA	CCAGAGAGAG	CTGAGCTGGA	
		GCGGGCCCG'	r ccagacaacc	TCCCGTCAA	CAGTTGGACT	GGTCTCTCTC	GACTCGACCT	
	61	AGACCCCACT	r gatggtgtgc	TGCCTTTCA	TCCAGGAAGA	AAGAAAGGAA	GGATTCTGAG	
	. _				AGGTCCTTCT	- • • • • • • • • • • • • • • • • • • •		
	121				TCTGTATACT			
					AGACATATGA			
	181				CGCGCCCTGA			
					GCGCGGGACT			
	241				TGGTTTGACG			
					ACCAAACTGC			
	301				ATTTGTTTCT			
	261				TAAACAAAGA			
	361				CAAAGCAAAA			
	421				GTTTCGTTTT			
	421				GGGAGTCGTG CCCTCAGCAC			
	481							
	401				CCACCTGCTC GGTGGACGAG			
	541						AGTGCAATAA	
	241				ACGTGACATG			
	601				GAAGGAAGGC			
	001				CTTCCTTCCG			
	661				TGCCTACGTG			
					ACGGATGCAC			
, ·	721				ATTTCCTCGA	· · - · · · · · · · · · · · · · · ·		
	•				TAAAGGAGCT			
	781	····			CATTGTCACT			** * *
					GTAACAGTGA			
	841	CATCCTTGAA	CTGTACAGGC	ACTGGGGCAA	AGAGGCCTTC	TGCCTCAGCC	ATCAGGAGGT	**********
		GTAGGAACTT	GACATGTCCG	TGACCCCGTT	TCTCCGGAAG	ACGGAGTCGG	TAGTCCTCCA /	
	901	TGCAACAGCC	AATCTTGCTT	GGGCTTCCGT	AGCAATACAC	CAGGTGCAGG	TGTGTAGAGG	
	·	ACGTTGTCGG	TTAGAACGAA	CCCGAAGGCA	TCGTTATGTG	GTCCACGTCC	ACACATCTCC	
	961				CCTACTCGAC			
		AGAGCGGTTT	GTCTGGCTTG	ACTTACTGAA	GGATGAGCTG	CATAGTTTCT	GTATGAAACT	
	1021			•	ACACATCATG			
					TGTGTAGTAC			
•	1081				GGACCACAAG		•	
				•	CCTGGTGTTC		•	
	1141				GAAGCCCATC			
					CTTCGGGTAG	• •		
	1201	•			TCAAGTGGCA			•
	1061		·		AGTTCACCGT	• • • •		
	1261				TAACAGGGAG			
	1221	·			ATTGTCCCTC			
					AGTGAGCCGA TCACTCGGCT			
					10A010GGC1	COLOGOROI		

	1381		AACAAGATCA				
		CGTCTACCAC	TTGTTCTAGT	CGCCATCGCG	GAAGAGGTTC	TGTCTGCTCT	TGTTGAAGTT
•••	1441		GTCTTCTGCG				
		CTACAAACGA	CAGAAGACGC	GTGACCGGAA	CGTGACACGA	TTGTACATGG	TGTCCTAGGC
	1501		TGCATCTACA				
		GGTGAGTCTT	ACGTAGATGT	CCCAATGGTA	CCTCTTCGAA	AGGATGGTGT	CGTAGACGTG
	1561		TGGCAAGGCC				
		GAGGCTCCTC	ACCGTTCCGG	AGTACGCGAA	GTTGGATGGT	CGTGCGTAGA	CGGCCCTGTA
	1621		CACTTTGACA				
		GCTCGATAAG	GTGAAACTGT	AACCAGGAAA	GCTCTTGTAC	ACCGGACCCT	AGAAACAGAT
	1681		CGGTCTTGTG				•
		GTACTAGGTA	GCCAGAACAC	CCTGTAGGAC	AAAACTTGAA	CTTTTTAACA	CGGCAAAATA
	1741		AAGAAGAACT				
		GTACAGACAC	TTCTTCTTGA	TAGCCGCCCA	AGGAATGGTG	TTGACCTTCG	TACGTCAGTG
	1801		TGCATGTATG				
		CCACCGTGTG	ACGTACATAC	GGTATGAAGT	TTTGTTGTTA	CCGGAGAAGT	GTCTGGAGCT
	1861		CTGCTAATTG				
			GACGATTAAC				
	1921		CAGAAGTTCG				
			GTCTTCAAGC				
	1981		TTCTCCCAGA	· ·			
			AAGAGGGTCT				
	2041		TCCAGCGAGT				
			AGGTCGCTCA				
	2101		GCCCTATACT				
			CGGGATATGA				
	2161		CTCCACAACC				
			GAGGTGTTGG				
	2221	-	TGCTCTGTGA				
			ACGAGACACT				
	2281		TTCTGGGCTG				
			AAGACCCGAC				
	2341						TCTACAATGC
			TCTCTGTTCG				
•	2401		CCCTGCTATA				
		· · · · · · · · · · · · · · · · · · ·			*		TCGGAGACGA
	2461						AAGAGACAGC
							TTCTCTGTCG
	2521						AGCTGAACGT
						· · · · · · · · · · · · · · · · · · ·	TCGACTTGCA
	2581						CGACTCAACC
						• • • · · · · · · · · · · · · · · · · ·	GCTGAGTTGG
							CAGAAGGTAC
		ACGAAGACAC	TGAAGCAAGA	AAAACAAAAG	TTCCCCACTT	TTGGGGGACA	GTCTTCCATG
	2701	CGTCGCATAT	CCATGTGAAG	CAGACGACTC	CCTGCTTGCC	GCACACACCT	CGGACAGTGA
	·	GCAGCGTATA	GGTACACTTC	GTCTGCTGAG	GGACGAACGG	CGTGTGTGGA	GCCTGTCACT
							•

	2761	GCAACCCAGG	CTCTGCCGTG	TTCAGACGTC	GGCTACTCCG	TGGCTCCACC	TGACCTCCGA
			GAGACGGCAC				
	2821	ATGCTATTTG	CTCCCAGGCC	AGCACTGCAC	TGTCTGGAGG	GGGCAGAGAC	CACAGGAGAG
		TACGATAAAC	GAGGGTCCGG	TCGTGACGTG	ACAGACCTCC	CCCGTCTCTG	GTGTCCTCTC
	2881	GTTCTTGCCT	GCATCCTCCC	ATGAGGGTGT	GGCCAGTTCC	CTAGTTCTGT	GCCATGCTGC
		CAAGAACGGA	CGTAGGAGGG	TACTCCCACA	CCGGTCAAGG	GATCAAGACA	CGGTACGACG
	2941	TGCTTGGTGG	CATTGGTTAG	GAATGGGACA	CACGCCCCTT	GTTGTGAAGT	TTACATGTGA
		ACGAACCACC	GTAACCAATC	CTTACCCTGT	GTGCGGGGAA	CAACACTTCA	AATGTACACT
	3001	CCTTCTTATA	GGTTAACTGA	GTTTGTGGCC	TGGGACACAT	GTAATGAAGG	TCACAGTCCA
		GGAAGAATAT	CCAATTGACT	CAAACACCGG	ACCCTGTGTA	CATTACTTCC	AGTGTCAGGT
	3061	CAGGTGACAG	AGAAATCCAA	ACTGTTGATT	ACAGGTGCAC	TACAGGTATG	CTCTTTCAGT
		GTCCACTGTC	TCTTTAGGTT	TGACAACTAA	TGTCCACGTG	ATGTCCATAC	GAGAAAGTCA
	3121	CTATCTGGGG	GCACATAGGT	GAGTCTGCTC	CACTCAGAAG	GAAGCATACC	TCTSCCCTCA
		GATAGACCCC	CGTGTATCCA	CTCAGACGAG	GTGAGTCTTC	CTTCGTATGG	AGASGGGAGT
	3181	TCCAGGGGAC	ACAGGGTACA	TCCCAGGCAT	CGGGGAACTG	AAGCTCTCAC	TTCAAACCAT
		AGGTCCCCTG	TGTCCCATGT	AGGGTCCGTA	GCCCCTTGAC	TTCGAGAGTG	AAGTTTGGTA
	3241	GTCAAAGAAT	TAAAACACCT	CCCCTCCCC	TCACTGTAGC	CTTCGGCAAC	TGCGCCAATC
	•	CAGTTTCTTA	ATTTTGTGGA	GGGGAGGGG	AGTGACATCG	GAAGCCGTTG	ACGCGGTTAG
	3301	CCTTTATACA	AAGAAAATAT	AAGTAAGGCA	TATAAATTTC	CTCCAGCAAG	CAAATCTTGT
		GGAAATATGT	TTCTTTTATA	TTCATTCCGT	ATATTTAAAG	GAGGTCGTTC	GTTTAGAACA
	3361	GGGTAAAAA	AAAAATGTG	AATTTTAACA	ACCTCTATAT	TTTCACTGTA	TGTTATGGCA
		CCCATTTTT	TTTTTTACAC	TTAAAATTGT	TGGAGATATA	AAAGTGACAT	ACAATACCGT
	3421		CACGTCCAAA				
							GATACGGACT
	3481		GCATGGCGTC				
			CGTACCGCAG				
	3541		GGAGGAGGTG				
						~~	AGGAGAGTGG
	3601		CCATAAGGAA				
			GGTATTCCTT			·- · · · · · · · · · · · · · · · · · ·	
	3661						CACAGGTTCA
			TCCTGTGGGG				
	3721	CACCGTGGGG	ATTTTCAGGA	TAGCATGGAG	ACCAGAGAAT	CCCGGTTCGG	TTGTTCTCAC
							AACAAGAGTG
	3781	TCGGTGAGCC	TTGAGAAGGA	AGAGACTGAC	CAGAAACACT	CACTCAGCAC	TCTGGCAGGA
							AGACCGTCCT
	3841						CAATACCATA
							GTTATGGTAT
	3901	CACACAGGAG	CTTGGCATTT	GCAAAGTCTA	TTCAGTTTCC	TTCCACACTC	TGACCCACGG
							ACTGGGTGCC
	3961	TTGTAGCGGA	GTGGGCTGAA	CACTGTAACA	CTGTACATGC	GATTTCCCCA	TGGGCTTCTA
							ACCCGAAGAT
	4021	AAATGTCACC	ATCTCCTCCC	CTGCTGTGTC	CTACTCCATT	TACTGGTTAC	AAGGTGATGT
							TTCCACTACA
٠							TGTATGCACA
		GTTGTTCTCT	TCGATAGTGT	TGTGGTCCCG	ACACGTGTGC	ACGTGTGTGT	ACATACGTGT

	4141	AGCACACAGA	TGTATGTACA	GCACACACAC	ACACACACAC	CCCAAAAGGA	GAGAAAAGGA
		TCGTGTGTCT	ACATACATGT	CGTGTGTGTG	TGTGTGTG	GGGTTTTCCT	CTCTTTTCCT
•••	4201				CCCATATTCA		
		TCTTTTGTAA	ATATTTTTCG	CTGTCGATGG	GGGTATAAGT	TTTTATCAAG	AAAAGGGACA
••	4261				TATCATGAGT		
							TAGTCACGTG
	4321				CACGTCTATA		
					GTGCAGATAT		
	4381				TCTTCTATCA		
					AGAAGATAGT		
	4441				TTCAGGCATT AAGTCCGTAA		
	4501				GGCATTGGGA		
	4501				CCGTAACCCT		
	4561			~	TCCTTGGATT		
	4561				AGGAACCTAA		
	4621				TGGTCAGTCT		
	4021				ACCAGTCAGA		
	4681				TGGGAAGACC		
					ACCCTTCTGG		
ij.	4741	ATGGGCCAAG	ACTCAGTTAG	GAGGCATAAA	TCTGCCCGGA	ATAAACTAGG	CCAGGATACA
		TACCCGGTTC	TGAGTCAATC	CTCCGTATTT	AGACGGGCCT	TATTTGATCC	GGTCCTATGT
	4801				ATTCACACAG		
					TAAGTGTGTC		
	4861				GGCATGGGAA		
					CCGTACCCTT		
	4921				CCAAAGGTTT		
					GGTTTCCAAA		= 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	4981				TATAAACAAT ATATTTGTTA		
					TTTTATCGCA	· ·—··	
	5041				AAAATAGCGT		
					GTTGTTTCTG		
	5101						GAACCTATAC
		TAAGTGCCAA					
	2161	ATTCACGGTT	ATGGTCACAC	TTCATAACTA	GGGCCCGTCG	TTTTATGTCG	GATTCCAAAC
	5221						CAGTGTAAAG
	J221	ATTTGTAGTT	AAGATAGAGT	CAAGTAGTCT	CCCGGACTCT	TCGACGCCCC	GTCACATTTC
	5281						CAATTGAATC
	3201	ATTTCATACG	ACCCGACCAC	CACCAGTCGG	AGGGGAACGG	TTCTTCTCTC	GTTAACTTAG
	5341	CTGTCCCCAG	CTCCCTCCAC	GCCTGAAGAG	TGACCAGTGC	TGGCCCGACG	GATCGCTGAG
		GACAGGGGTC	GAGGGAGGTG	CGGACTTCTC	ACTGGTCACG	ACCGGGCTGC	CTAGCGACTC
	5401	ATATTCTCCC	ATAATGGCAA	AAAAATAGGC	AGTTTGATGT	GACCTGTTTA	GTGTGGCTCT
		TATAAGAGGG	TATTACCGTT	TTTTTATCCG	TCAAACTACA	CTGGACAAAT	CACACCGAGA
	5461	CCTCTTTTGA	GCATGTGTTA	GCATTTTTAT	TTTATACTCA	TCCAGTGAAC	TCTGCTCTTC
		GGAGAAAACT	CGTACACAAT	CGTAAAAATA	AAATATGAGT	AGGTCACTTG	AGACGAGAAG

Figure 19 (con't) PDE10A compiled

				•				
	5521	CAAGTGTGTT	CATGTATGTG	CTAGATATAT	TAGCACAGCC	TGCCTTCTGC	TGCACAACGC	
•			GTACATACAC					:
	5581	CTTAGAGACC	CGGCCTTTCA	ATGAGCTTAG	CTTGTGCTCT	GTTTCTGCTC	TCTTAGGTCT	
			GCCGGAAAGT					
	5641	AAACTATGGT	GTCAGTTTTA	ATAGAACAAA	AGTATGCATC	TTGCCTTGGC	TTGAGCCTTT	
		TTTGATACCA	CAGTCAAAAT	TATCTTGTTT	TCATACGTAG	AACGGAACCG	AACTCGGAAA	
	5701		TGCTGACTTC					
		AGCAAAAGTT	ACGACTGAAG	AGGGGAAAGA	GAGGACACGA	GTGGAATGGA	AAGGTCTCAC	
	5761	TAAGGGACAA	CTTTTAAGGA	GGCGTGTCCC	TGGTAGGGGC	ATCCCTGTTC	ACCAGGTGCC	
		ATTCCCTGTT	GAAAATTCCT	CCGCACAGGG	ACCATCCCCG	TAGGGACAAG	TGGTCCACGG	
	5821	TGTCATCACC	CCACTTGACT	GACATCTACC	CTGGTGACTA	TGGGTTCCTC	TTGTTTGTAG	
							AACAAACATC	
	5881	GGAACGGTGG	CTCCAGGTGG	AGGCATCAAT	CTGTTGGGTT	CTGGTTCCCG	GCTGCCTTTG	
		CCTTGCCACC	GAGGTCCACC	TCCGTAGTTA	GACAACCCAA	GACCAAGGGC	CGACGGAAAC	
	5941	GTTTTGAAAG	TCTCTTCTCT	GTATATTCCT	ACCCTGCATT	TGCTTTGTGT	GGTGCTGATG	
		CAAAACTTTC	AGAGAAGAGA	CATATAAGGA	TGGGACGTAA	ACGAAACACA	CCACGACTAC	
	6001	CTGTGGCAGT	AGGATCTTGG	ATGACTCTCC	ATCAGTCACA	GACTCCCCCT	GTTGCAAAGT	
	-	GACACCGTCA	TCCTAGAACC	TACTGAGAGG	TAGTCAGTGT	CTGAGGGGGA	CAACGTTTCA	
	6061	GTCAGGCTGA	CTCGACAGTC	ACCGTAAAAT	CTGAGTCAGT	CACACACAGG	CTGTCAGCCA	•
	.000_	CAGTCCGACT	GAGCTGTCAG	TGGCATTTTA	GACTCAGTCA	GTGTGTGTCC	GACAGTCGGT	
	6121	CGGCTTCCAC	TTGCATGGCT	ATTCTATTTT	CACACGTGAG	TTTCTGTTGC	TGGCTGGCTG	
		GCCGAAGGTG	AACGTACCGA	TAAGATAAAA	GTGTGCACTC	AAAGACAACG	ACCGACCGAC	
	6181						CATCATTCTC	
	0202	TGACCGTAAT	AGATACGATT	CAACTTTAGT	CCTCACACGG	GTCGTCTCGG	GTAGTAAGAG	
	6241						CATTTCATGC	
	0212	TGACAGAAAC	TTTGTTTCGA	CATGCCAAAC	TAGCTACTTG	CATAAATTTC	GTAAAGTACG	
	6301	AATGACAAAG	TGCTCAGTAG	TGGAAGGCAG	GCTGTGACCA	GTCTGCCTGC	TCCTTACTAT	•
		TTACTGTTTC	ACGAGTCATC	ACCTTCCGTC	CGACACTGGT	CAGACGGACG	AGGAATGATA	
	6361	AATTGTGAGG	ATTTGTTACT	GGAACAGTAC	ATGGAGGCCT	GACCTTGTGG	GGGCACAGGG	
		TTAACACTCC	TAAACAATGA	CCTTGTCATG	TACCTCCGGA	CTGGAACACC	CCCGTGTCCC	
	6421	TGGAACCTTA	GCTGAATATA	GTGTGTGTCT	CAAGAGGAAG	TCAGGGTACT	AGCTCAGTGC	
		ACCTTGGAAT	CGACTTATAT	CACACACAGA	GTTCTCCTTC	AGTCCCATGA	TCGAGTCACG	
	6481	TCAATCTCCA	GGTACTATAT	ATACATTTGC	CCGTTTTATC	TCTAATGTGA	AATAAATCCC	
		AGTTAGAGGT	CCATGATATA	TATGTAAACG	GGCAAAATAG	AGATTACACT	TTATTTAGGG	
-	6541	CAAACACTTG	TTTATCGTGT	AGCGTACCTA	AAAGACTATT	CTATTATGGG	TGTCCCCACT	
•		GTTTGTGAAC	AAATAGCACA	TCGCATGGAT	TTTCTGATAA	GATAATACCC	ACAGGGGTGA	
·	6601	TTCTTGGTTT	GGTCACCCCG	ATCCCCCGGT	CTTCTGCTGT	ATCTAGAACA	GTGACTATAA	
		AAGAACCAAA	CCAGTGGGGC	TAGGGGGCCA	GAAGACGACA	TAGATCTTGT	CACTGATATT	•
	6661	ATGATGTATG	GGAATAGTGT	TTCCATATGA	TCTGTTGTCT	GGAGTATATG	CTACATGTTC	
		TACTACATAC	CCTTATCACA	AAGGTATACT	AGACAACAGA	CCTCATATAC	GATGTACAAG	
	6721	ATTTACTGTA	CAAAAACCCA	GTGCAGCTGA	TGATGCAAAG	CAGTCTCTCT	CTGTGTACAG	:
		TAAATGACAT	GTTTTTGGGT	CACGTCGACT	ACTACGTTTC	GTCAGAGAGA	GACACATGTC	
	6781	TGCCCCACCT	ATTTAAAAAT	CACGTACAAN	CCCAGAACAC	TGTGAAACAC	TTAACATAAG	
		ACGGGGTGGA	TAAATTTTTA	GTGCATGTTN	GGGTCTTGTG	ACACTTTGTG	AATTGTATTC	
	6841	AAACAAACGC	AGCGTCTGGA	TTCTTTCCAA	GGAGAGCAGC	TTTCTCCACA	GGAACACAGT	
	- -	TTTGTTTGCG	TCGCAGACCT	AAGAAAGGTT	CCTCTCGTCG	AAAGAGGTGT	CCTTGTGTCA	
		· — · —						

6901	AACAAAAGAG	GTCCGCCGCC	ATCCACACCC	AGCCAAGACA	CCTCAGAGGC	CATAGGGACA	
 	TTGTTTTCTC	CAGGCGGCGG	TAGGTGTGGG	TCGGTTCTGT	GGAGTCTCCG	GTATCCCTGT	
6961	ACCTCCTTGC	TGGCCAACAC	CTGCTGGAGC	AGGGCACAGG	TCCCAGCAAC	TGATCCTCAG	
 	TGGAGGAACG	ACCGGTTGTG	GACGACCTCG	TCCCGTGTCC	AGGGTCGTTG	ACTAGGAGTC	
7021	TGGATGGGTC	CGCAGTCAAA	GCCTTAATGG	GCTCTCTTTT	GAAGGGGAAA	GAAANNTTTC	
 	ACCTACCCAG	GCGTCAGTTT	CGGAATTACC	CGAGAGAAAA	CTTCCCCTTT	CTTTNNAAAG	
7081	AAGCTTATGA	TATCCAACAT	TATTATAGTT	GATGAGTTAG	TAAATTCCGA	AAAAAAAGA	• • • •
 	TTCGAATACT	ATAGGTTGTA	ATAATATCAA	CTACTCAATC	ATTTAAGGCT	TTTTTTTTCT	
7141	TGATTTTATA	TGTATGACAT	AAAAAAAATC	TTTGTAAAGT	GCGCAAGTGC	AATAATTTAA	
 	ACTAAAATAT	ACATACTGTA	TTTTTTTAG	AAACATTTCA	CGCGTTCACG	TTAAAATT	
7201	AGAGGTCTTA	TCTTTGCATT	TATAAATTAT	AAATATTGTA	CATGTGTGTA	ATTTTTCATG	
 	TCTCCAGAAT	AGAAACGTAA	ATATTTAATA	TTTATAACAT	GTACACACAT	TAAAAAGTAC	
7261	TATTCATTTG	CAGTCTTTGT	ATTTAAAAAA	ACTTTACTGT	TATGTTTGTA	TAATAGAACA	
 	ATAAGTAAAC	GTCAGAAACA	TAAATTTTTT	TGAAATGACA	ATACAAACAT	ATTATCTTGT	
7321	TTAATCATTT	ATTATAACTC	AGACAAGGTG	TAAATAAATT	CATAATTCAA	ACAGCCAGTA	
 	AATTAGTAAA	TAATATTGAG	TCTGTTCCAC	ATTTATTTAA	GTATTAAGTT	TGTCGGTCAT	
7381	TATATGCATA	TATGGGTGTT	ACATTGCAAA	AATCTCTATC	TTTGTTCTAT	TCACATGCTT	
 	ATATACGTAT	ATACCCACAA	TGTAACGTTT	TTAGAGATAG	AAACAAGATA	AGTGTACGAA	
 7441	AAAGAAGTAA	GAAATCTTTT	GTGGATATGT	AATTATACAT	ATAAAGTATA	TATATATGTA	
 	TTTCTTCATT	CTTTAGAAAA	CACCTATACA	TTAATATGTA	TATTTCATAT	ATATATACAT	
7501	TGATACATGA	AATATATTTA	GAAATGTTCA	TAATTTTAAT	GGATATTCTT	TGGTGTGAAT	
 . 4	ACTATGTACT	TTATATAAAT	CTTTACAAGT	ATTAAAATTA	CCTATAAGAA	ACCACACTTA	
7561	AATTGAATAC	AACATTTTTA	AAATGAAAAA	AAAAAAAAA	АААААААА	AAAAAAA	
 ·	TTAACTTATG	TTGTAAAAAT	TTTACTTTTT	TTTTTTTTT	TTTTTTTTT	TTTTTTT	